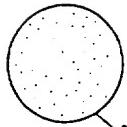
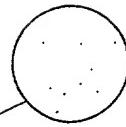


FIG. 1

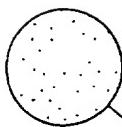


[ SEQ. ID NO: 3]

X-C-C-T-T-G-A-G-A-T-T-T-C-C-C-T-C  
5' 3'



G-G-A-A-C-T-C-T-A-A-A-G-G-G-A-G-X  
3' 5'  
[ SEQ. ID NO: 4]



X-C-C-T-T-G-A-G-A-T-T-T-C-C-C-T-C  
G-G-A-A-C-T-C-T-A-A-A-G-G-G-A-G-X

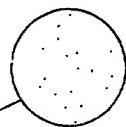


FIG.2

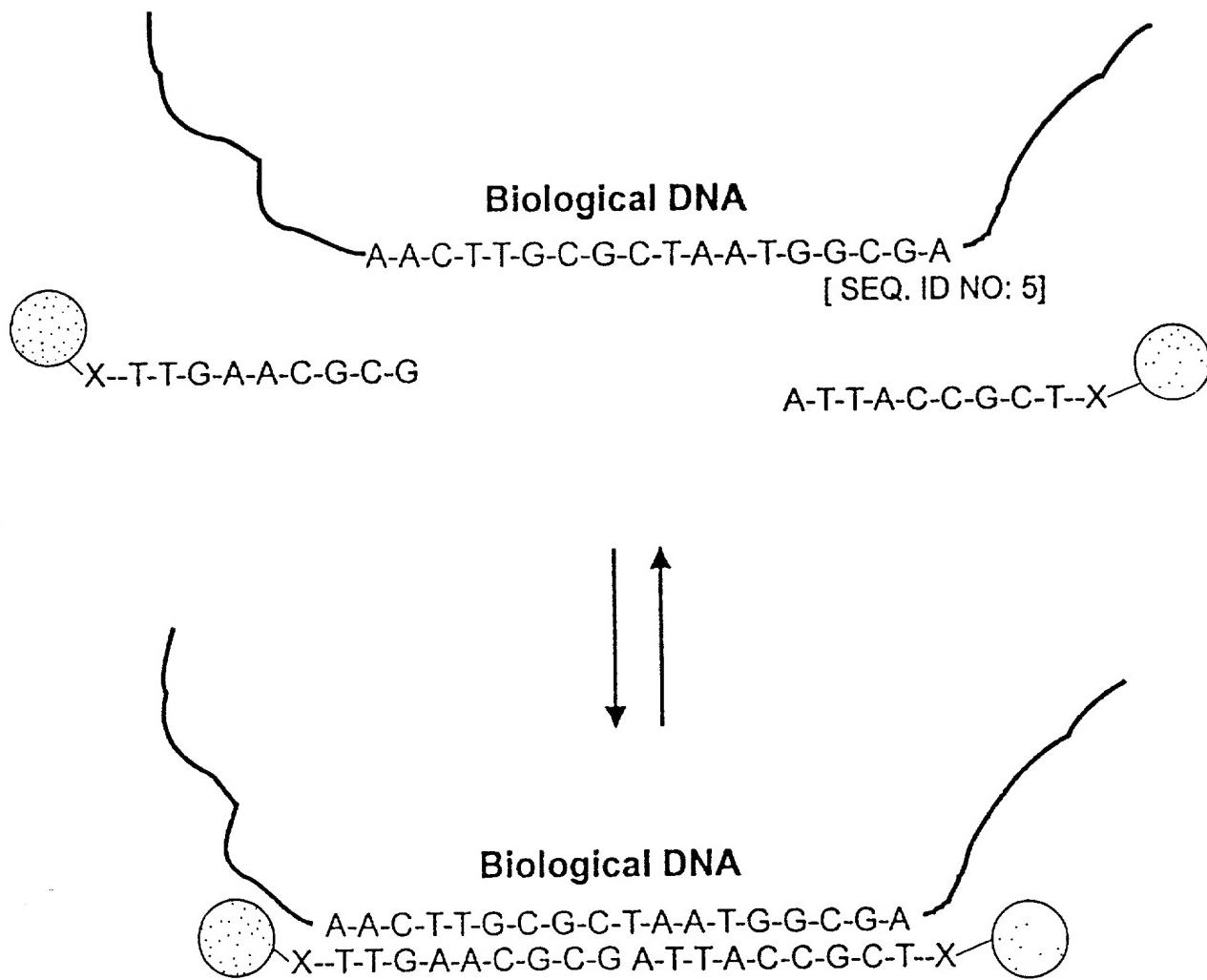
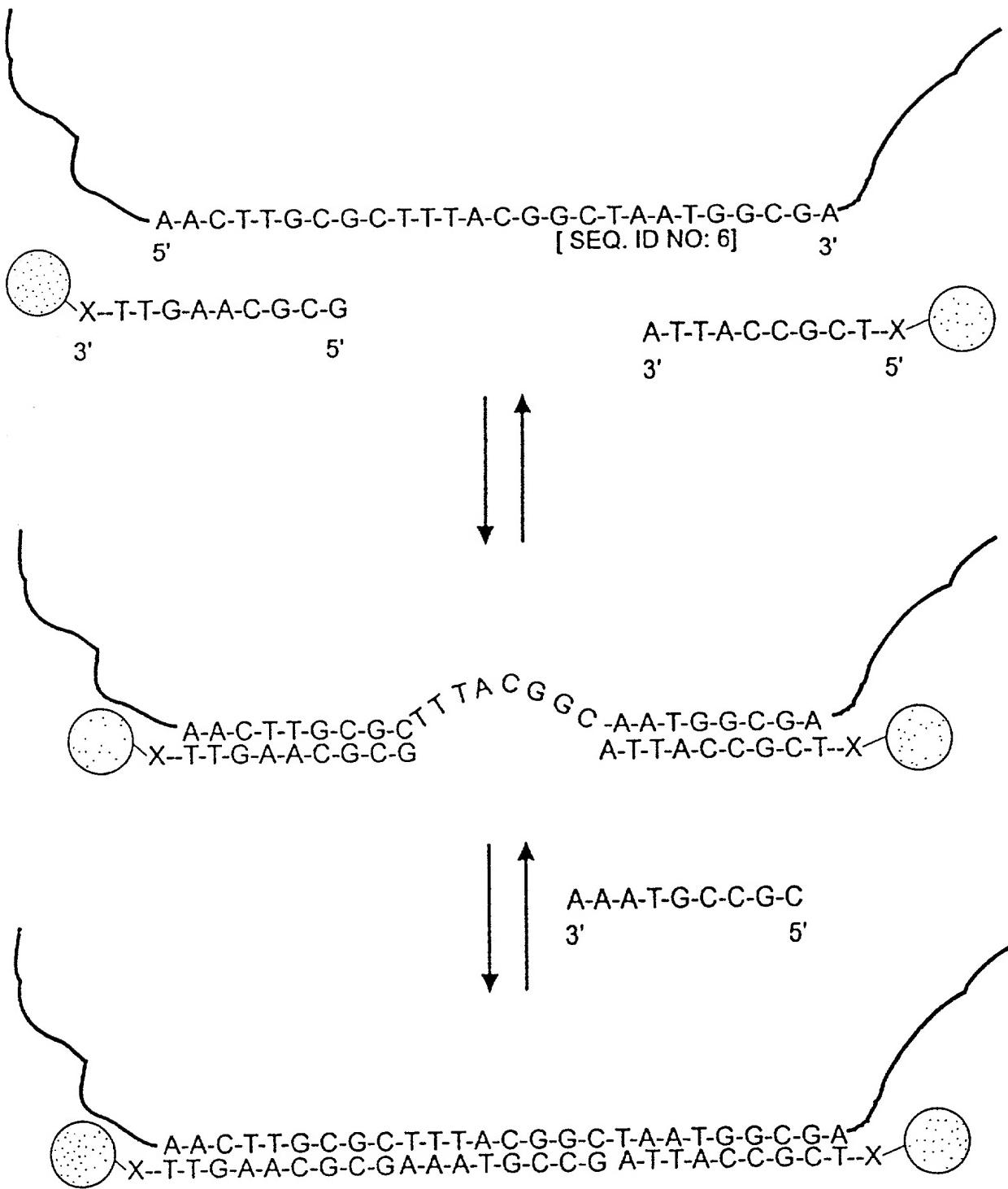


FIG.3

T020300 - 2023-02-06 00



**FIG. 4**

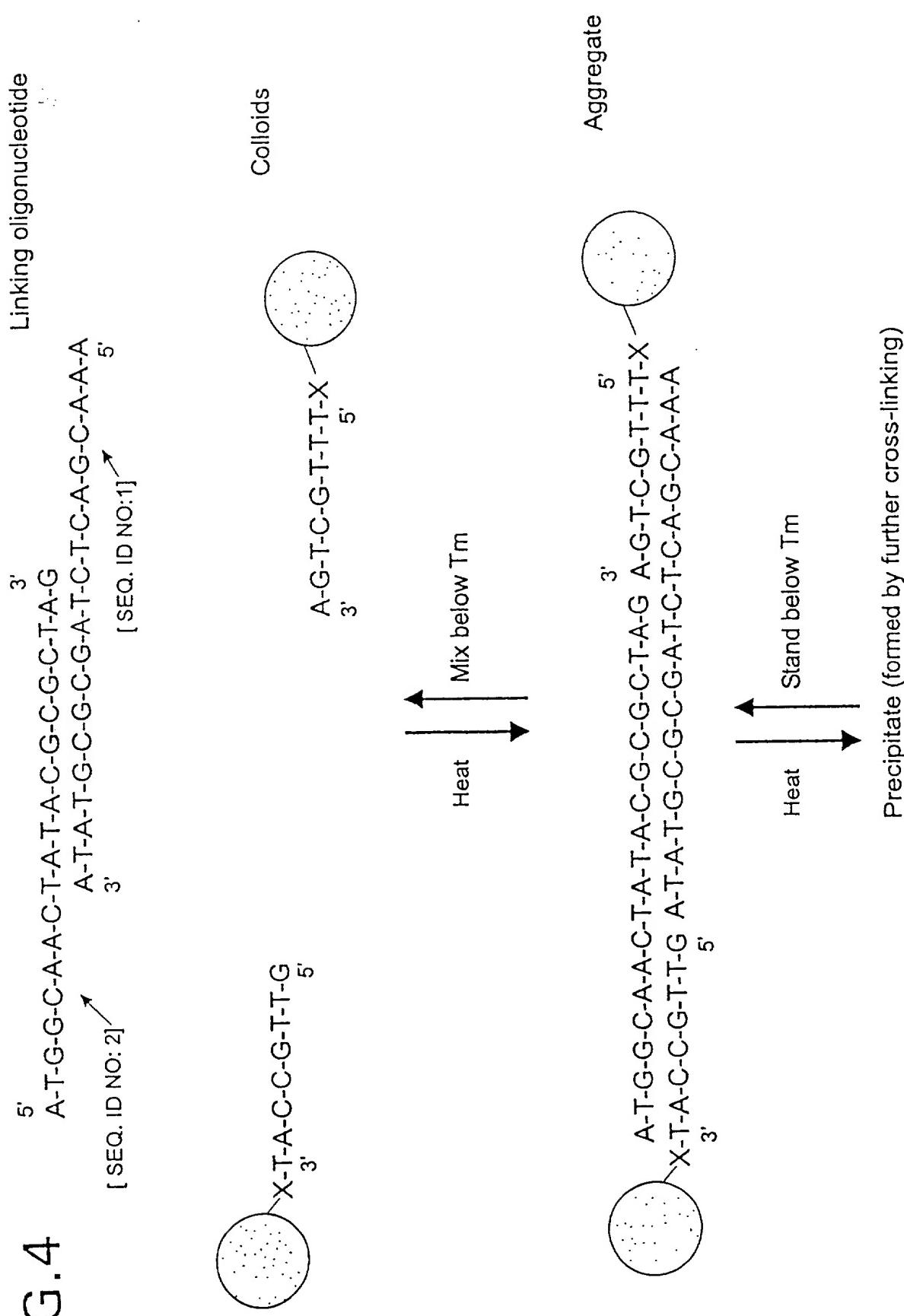
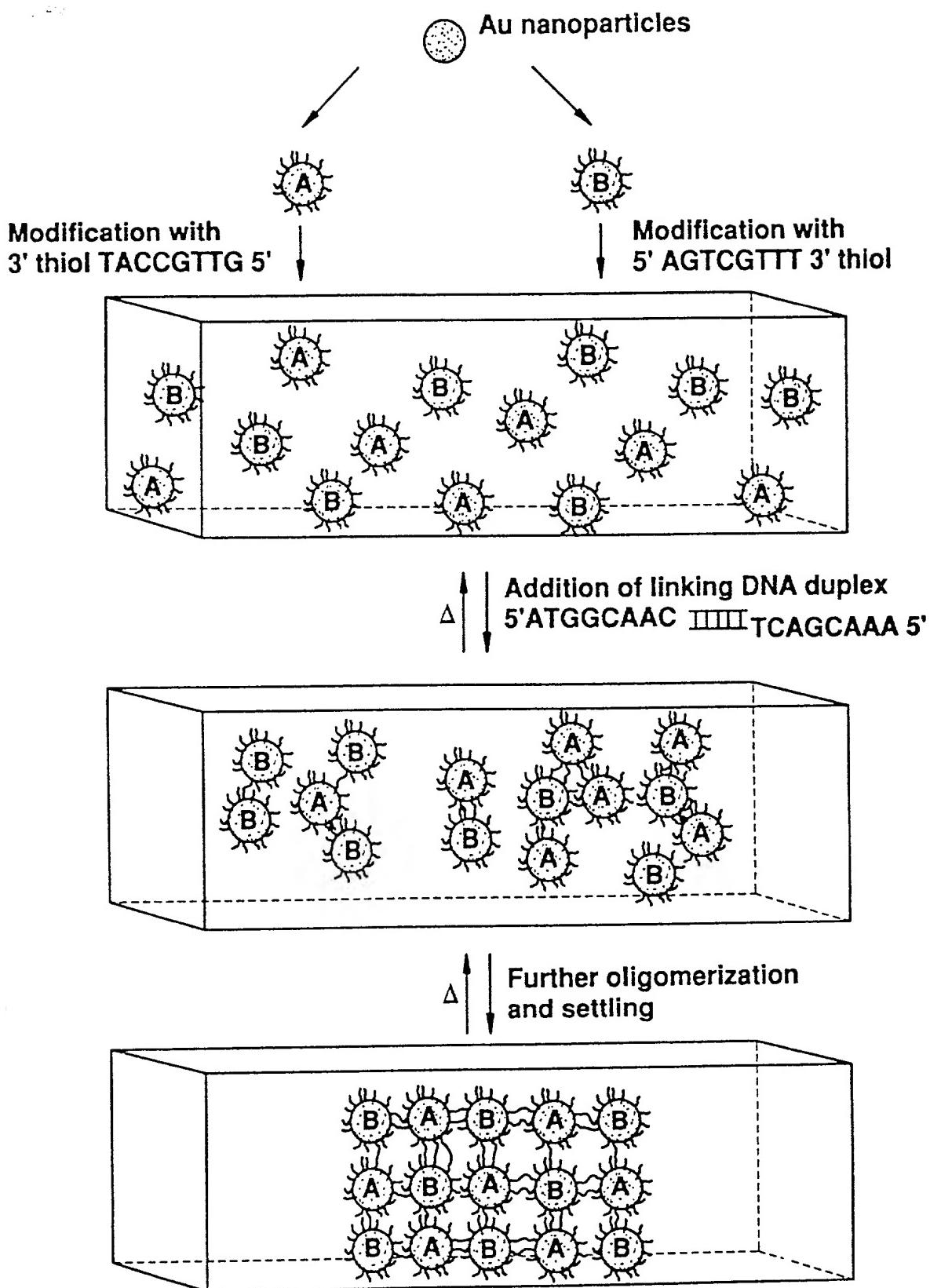


FIG.5



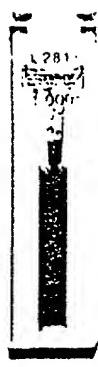


FIG.6A



FIG.6C

FIG. 7

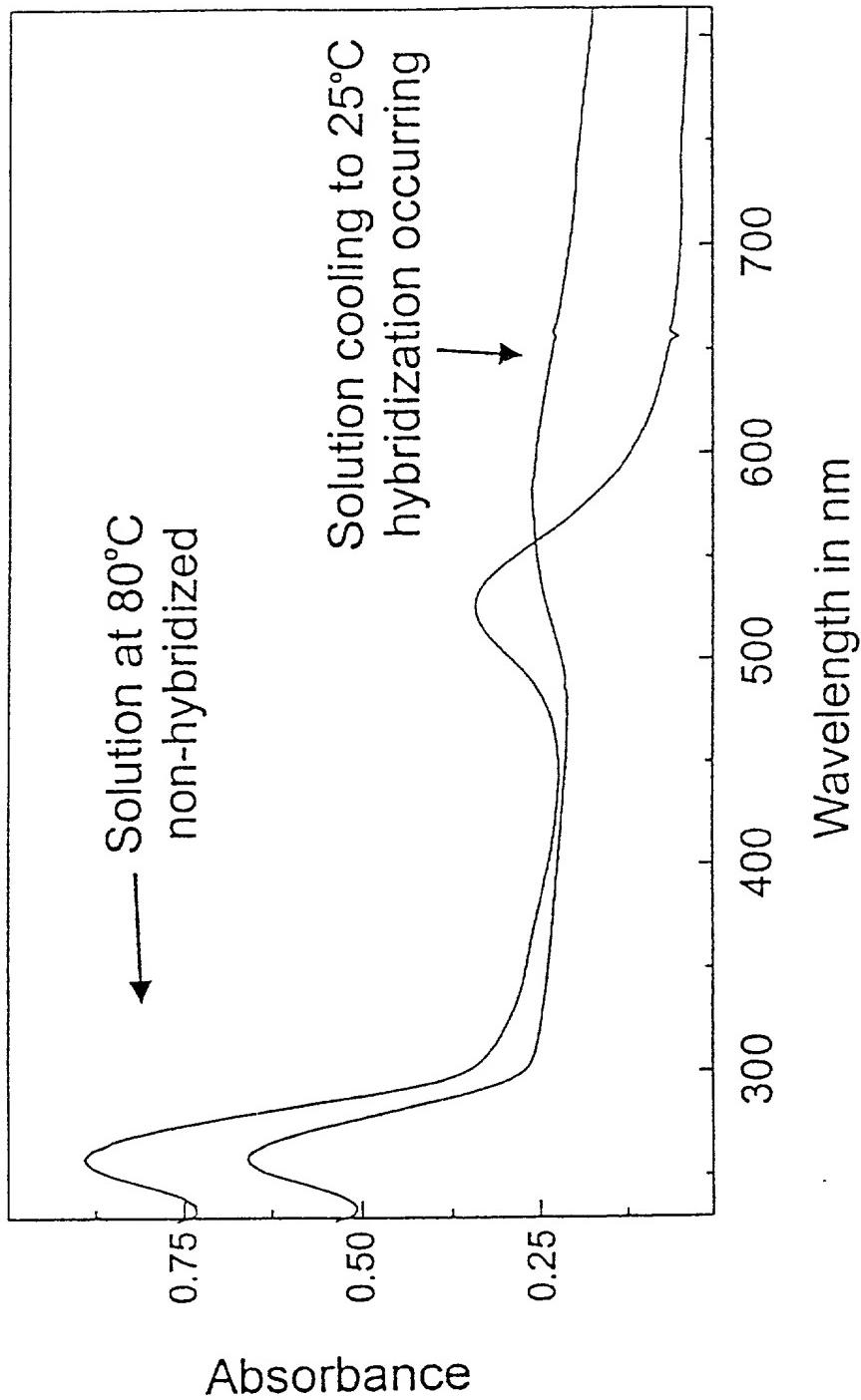


FIG. 8B

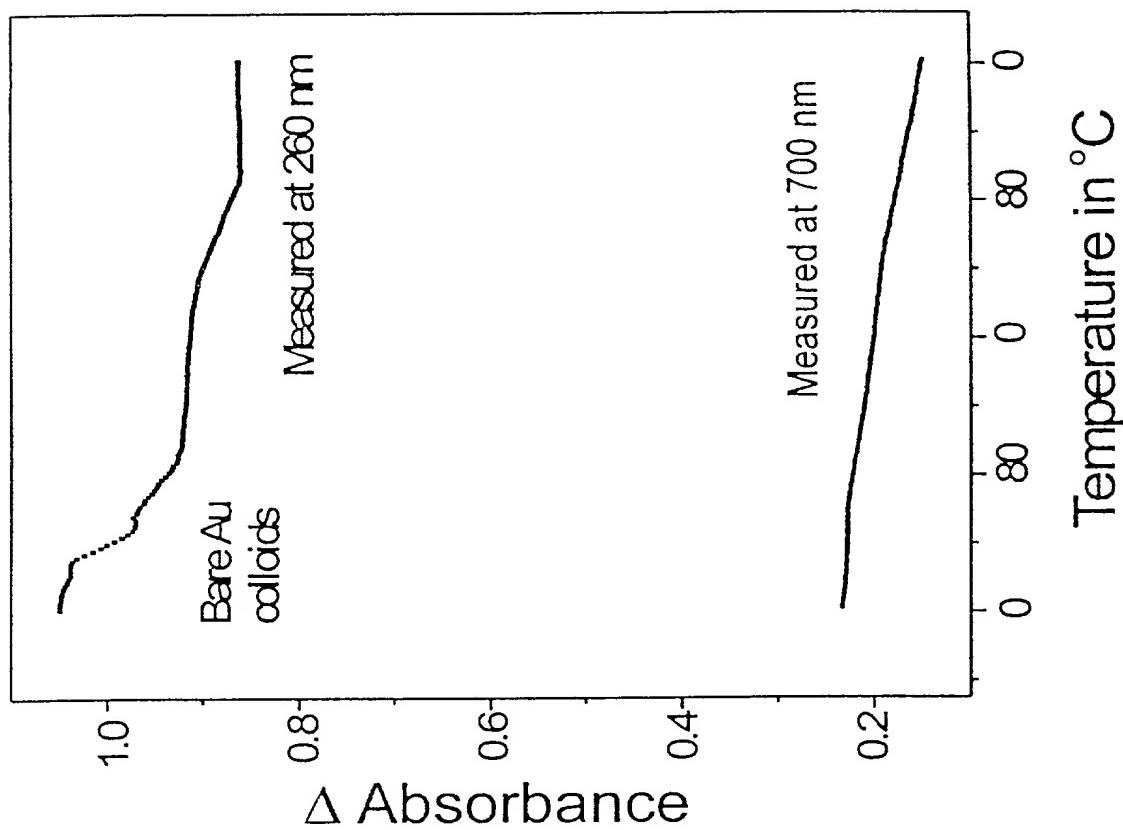
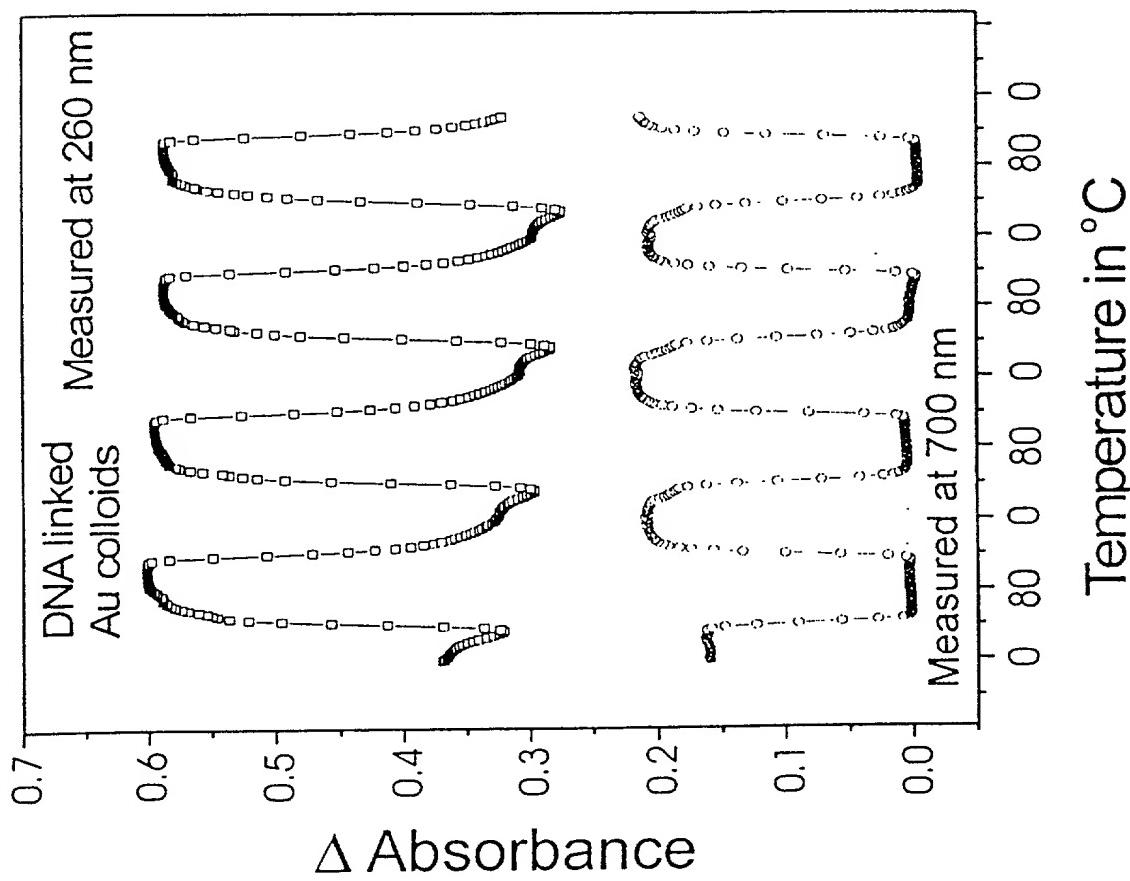


FIG. 8A



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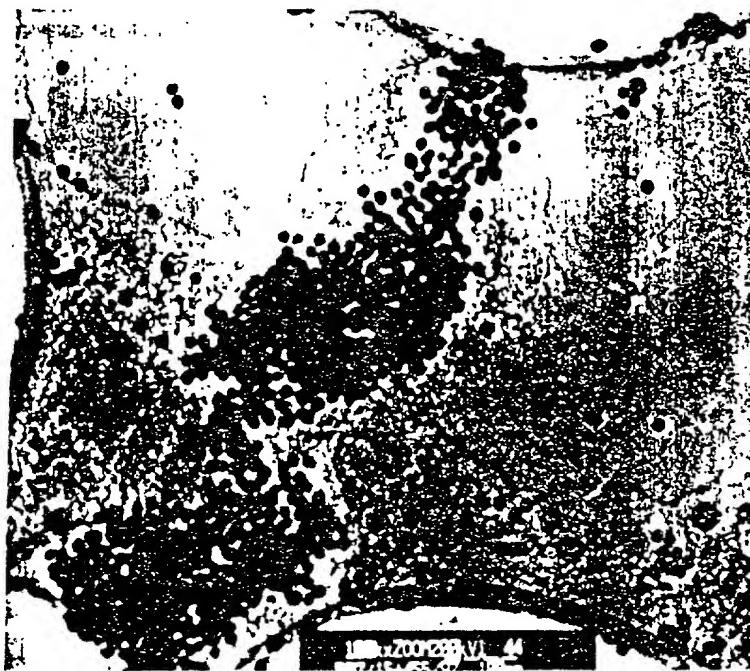


FIG.9A

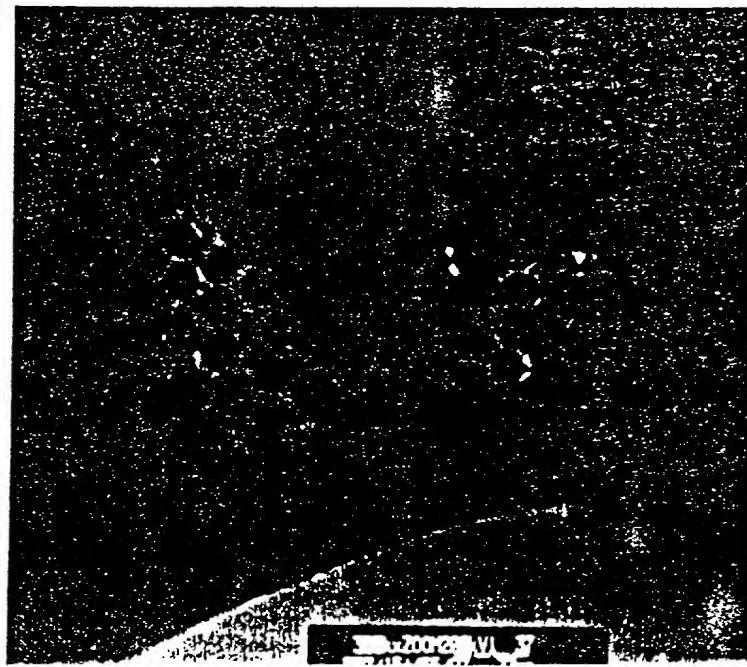
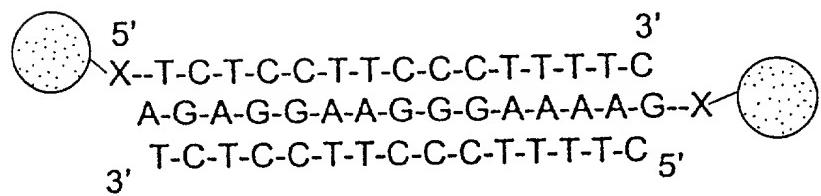
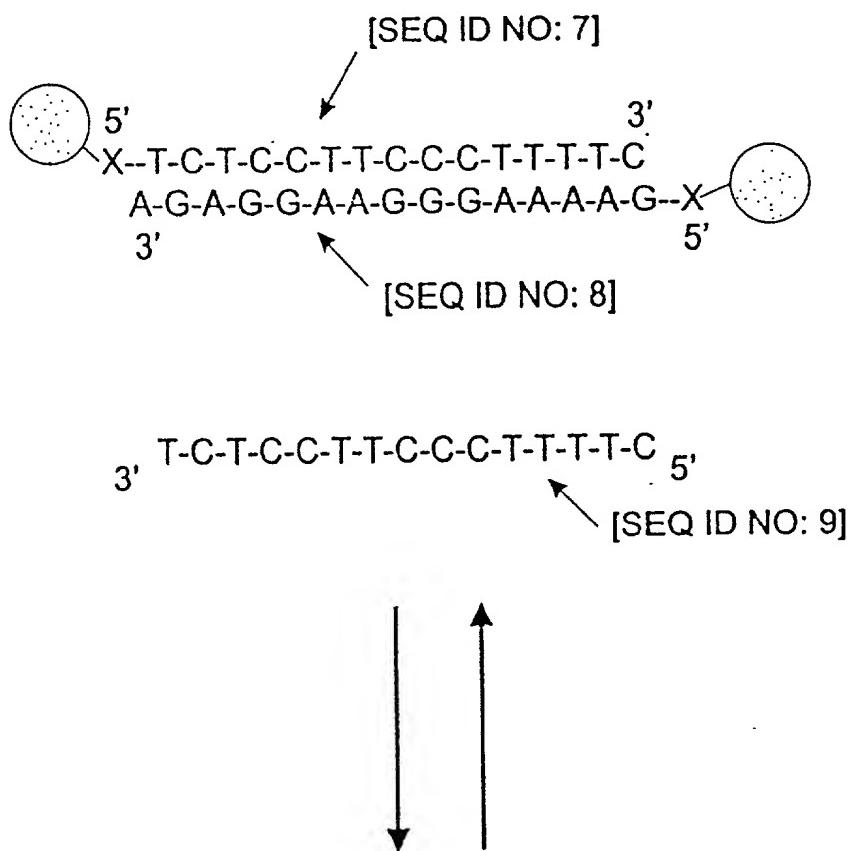
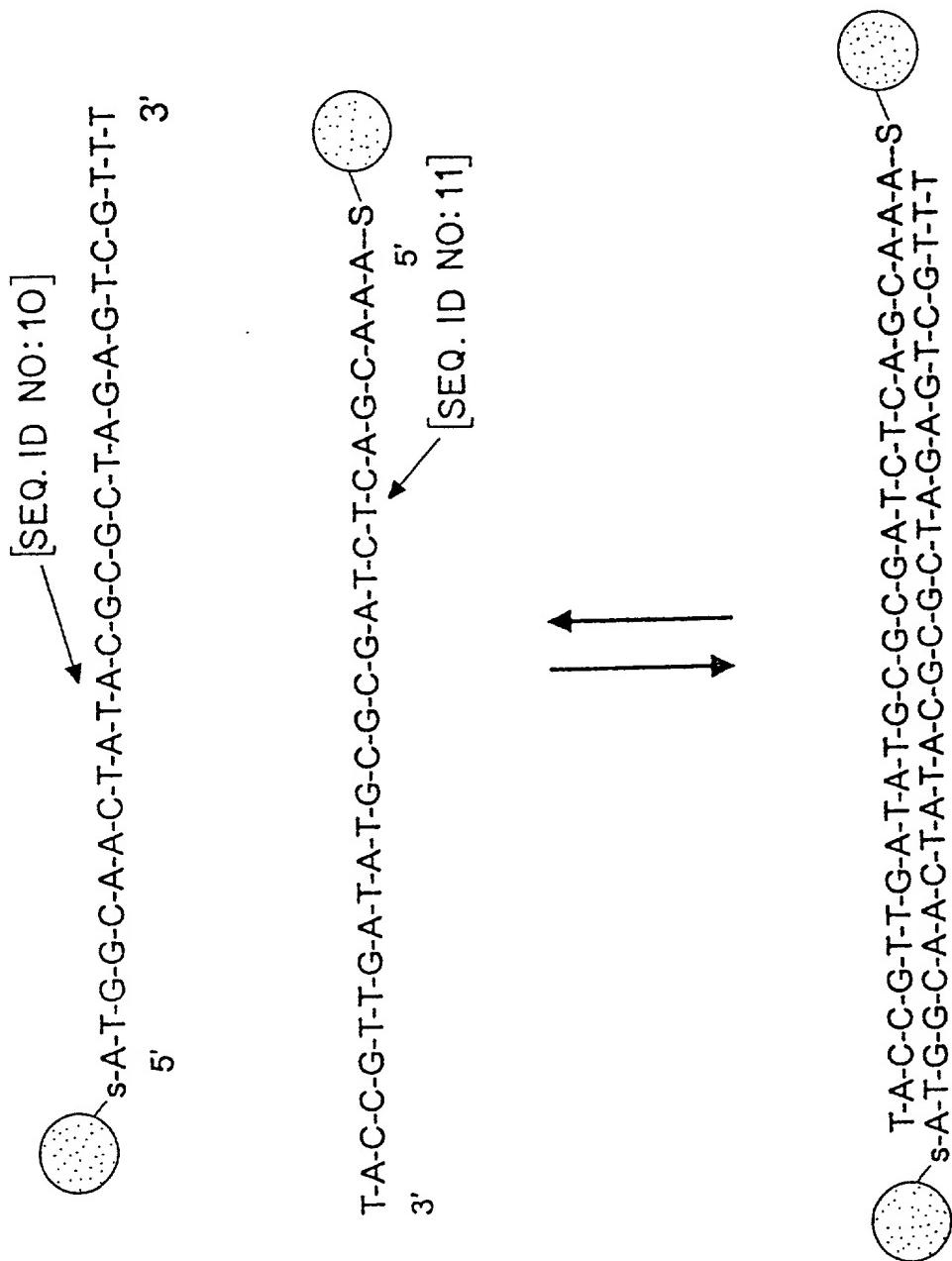


FIG.9B

FIG.10



## FIG. 11



**FIG. 12A**Complementary Target

[SEQ. ID NO:12]

1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C  
 5' A-G-C-A-T-G-G-T-C-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C

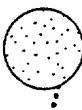
3

[SEQ. ID NO:14]

2

T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

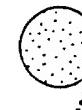
[SEQ. ID NO:13]

**FIG. 12B**Probes without Target1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C

2

T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

4**FIG. 12C**Half Complementary Target1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C  
 5' A-G-C-A-T-G-G-T-C-G-A-T-A-G-G-A-T-G-G-C-A-A-C-T-A-T-A-C-G-C

2

T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

[SEQ. ID NO:15]

4**FIG. 12D**Target - 6 bp1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C  
 5' G-T-C-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C

2

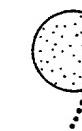
T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

5**FIG. 12E**One bp Mismatch1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C  
 5' A-G-C-A-T-G-G-T-T-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C

2

T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

6

[SEQ. ID NO:17]

**FIG. 12F**Two bp Mismatch1

3' T-C-G-T-A-C-C-C-A-G-C-T-A-T-C-C  
 5' A-G-C-A-T-G-T-T-G-A-T-A-G-G-A-A-A-C-G-A-C-T-C-T-A-G-C-G-C

2

T-T-T-G-C-T-G-A-G-A-T-C-G-C-G

7

[SEQ. ID NO:18]

FIG.13A

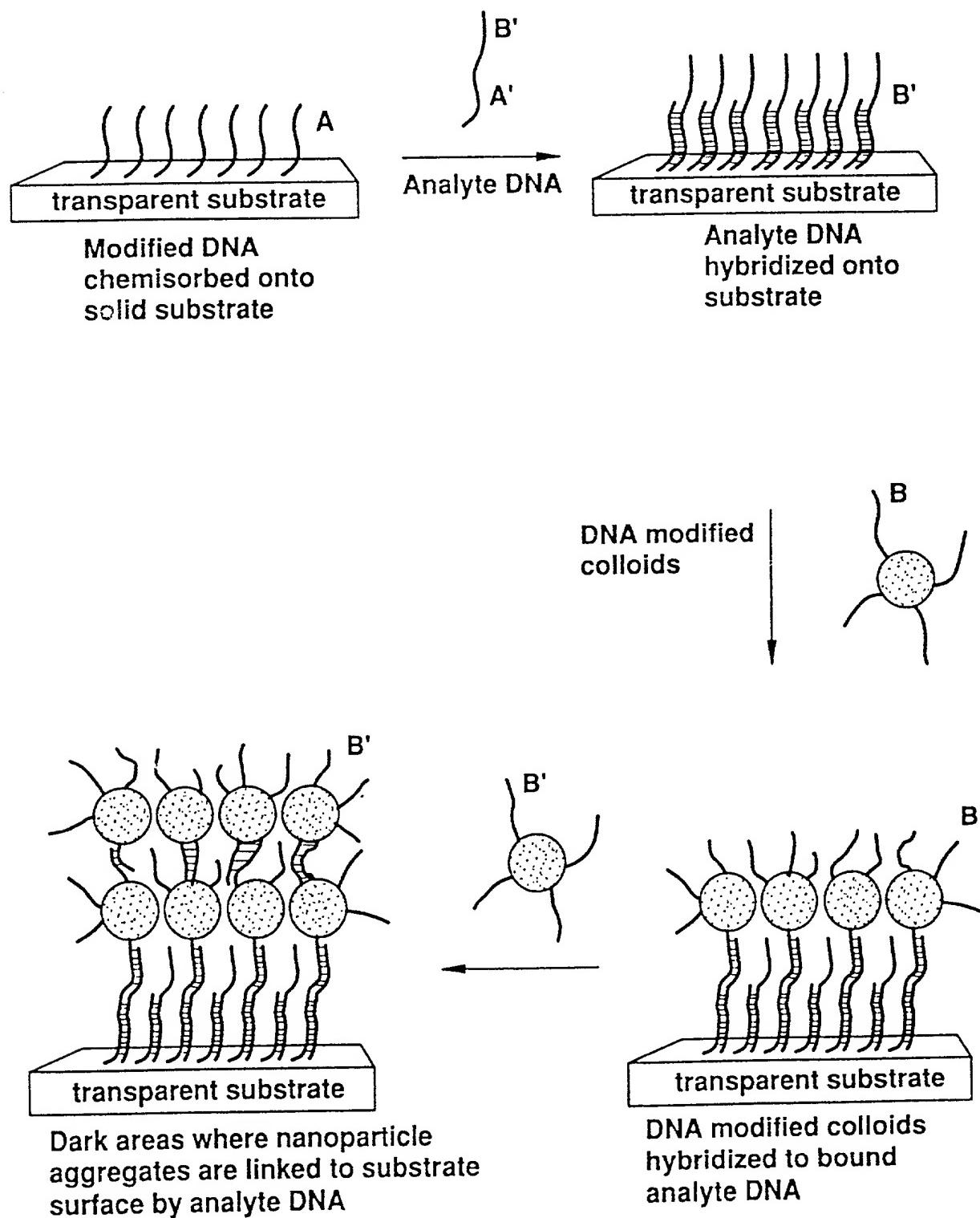


FIG. 13B

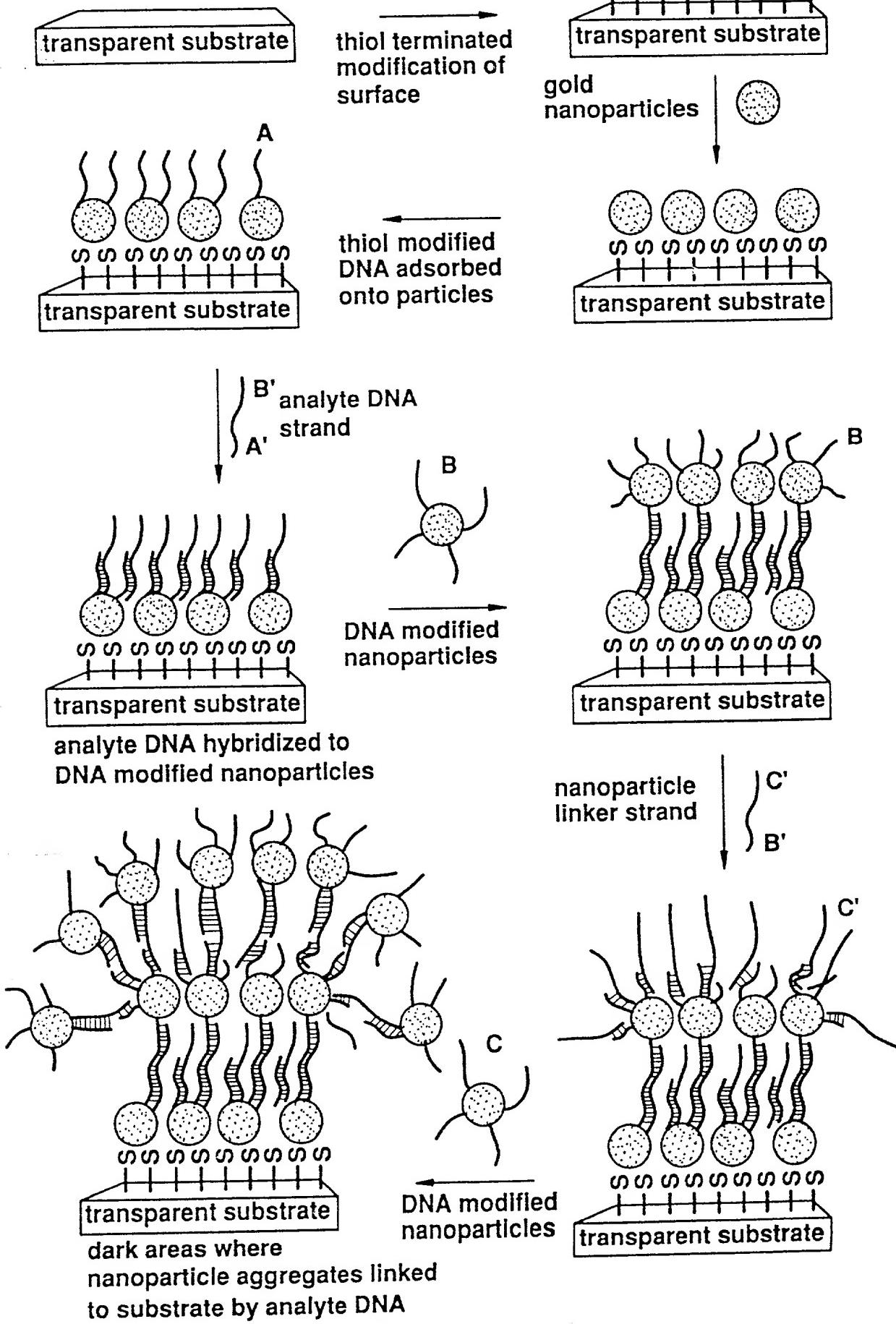


FIG.14A

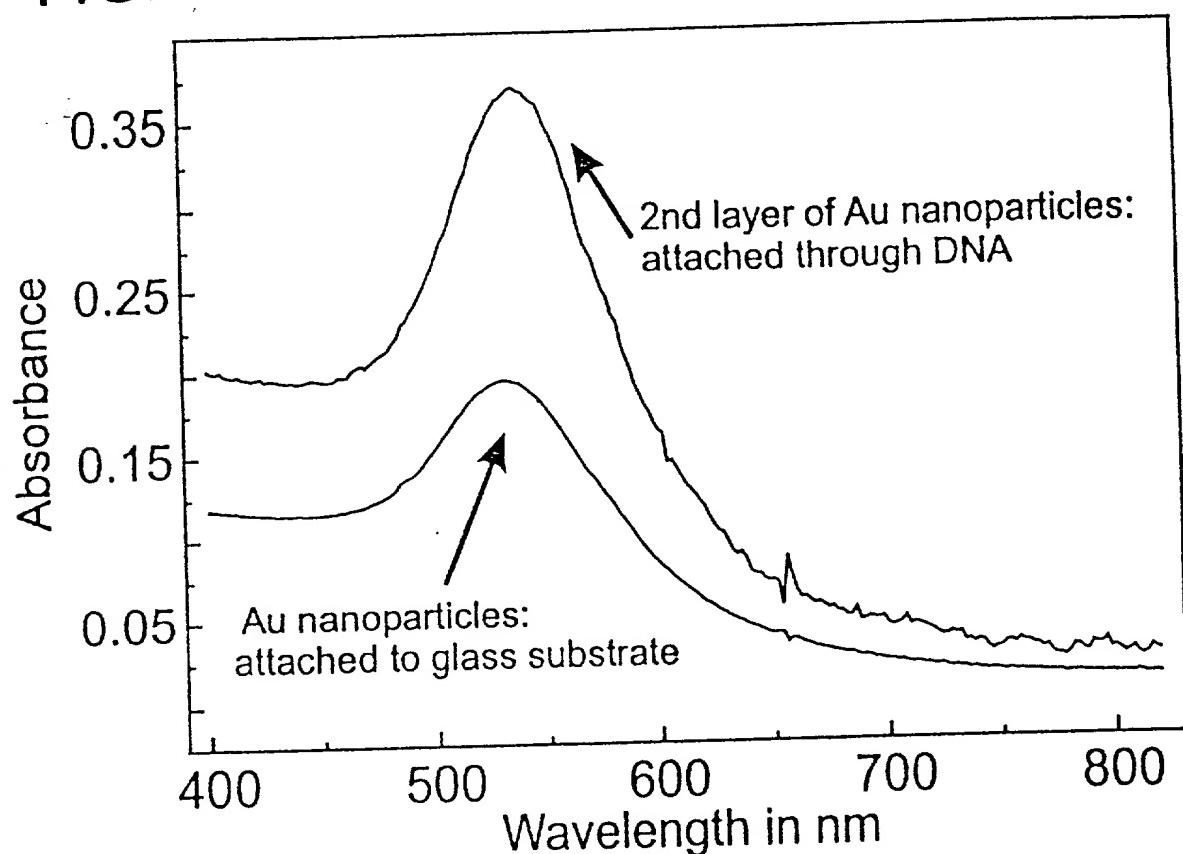
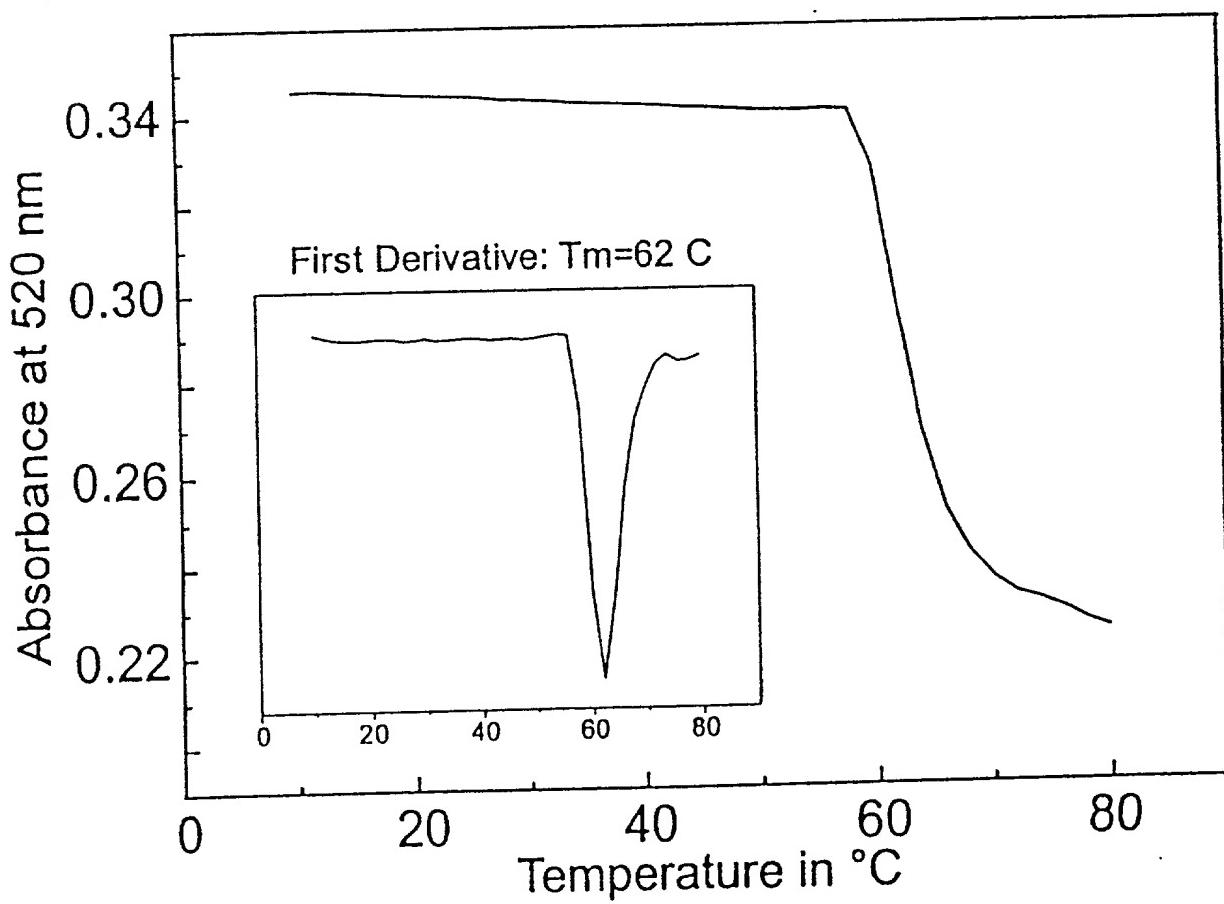


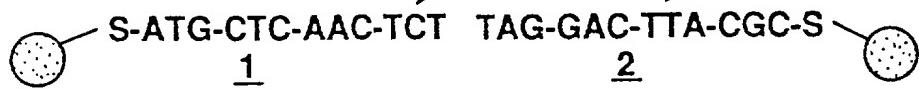
FIG.14B



**FIG15A**Probes with No Target

SEQ ID NO:19

SEQ ID NO:20

**FIG15B**Half-Complementary Target3

SEQ ID NO:21

5' TAC-GAG-TTG-AGA-GAG-TGC-CCA-CAT 3'

**FIG15C**Complementary Target

Tm=53.5°C

4

SEQ ID NO:22

5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCG 3'

**FIG15D**ONE Base-Pair Mismatch at Probe Head

Tm=50.4°C

5

SEQ ID NO:23

5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCT 3'

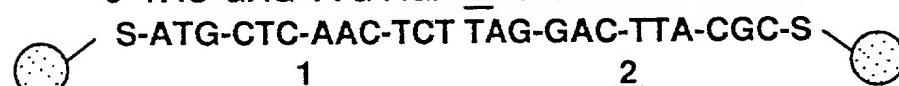
**FIG15E**ONE Base-Pair Mismatch at Probe Tail

Tm=46.2°C

6

SEQ ID NO:24

5' TAC-GAG-TTG-AGA-CTC-CTG-AAT-GCG 3'

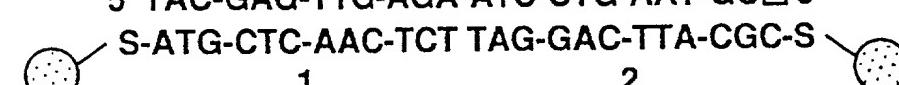
**FIG15F**ONE Base Deletion

Tm=51.6°C

7

SEQ ID NO:25

5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GC□ 3'

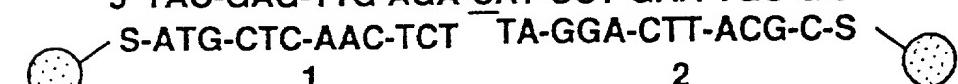
**FIG15G**ONE Base-Pair Insertion

Tm=50.2°C

8

SEQ ID NO:26

5' TAC-GAG-TTG-AGA-CAT-CCT-GAA-TGC-G 3'



**FIG. 16A**

24 Base Template

5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCG 3'  
—S-ATG-CTC-AAC-TCT TAG-GAC-TTA-CGC-S —  
1  
2

**FIG. 16B**

48 Base Template with Complementary 24 Base Filler

5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-ATC-CTG-AAT-GCG 3'  
—S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT TAG-GAC-TTA-CGC-S —  
1  
2

**FIG. 16C**

72 Base Template with Complementary 48 Base Filler

5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-TAT-AT—GGA-CGG-TTT-ACG-GAC-AAC-ATC-CTG-AAT-GCG 3'  
—S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT-ATA-TAA-CCT-GCG-AAA-TGC-CTG-TTG TAG-GAC-TTA-CGC-S —  
1  
2

FIG. 17A

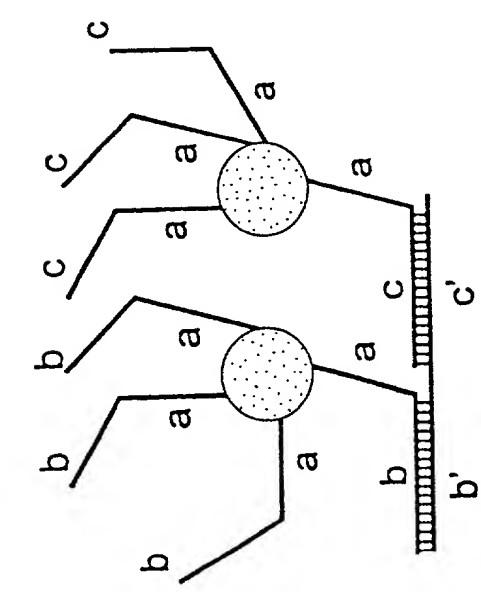


FIG. 17B

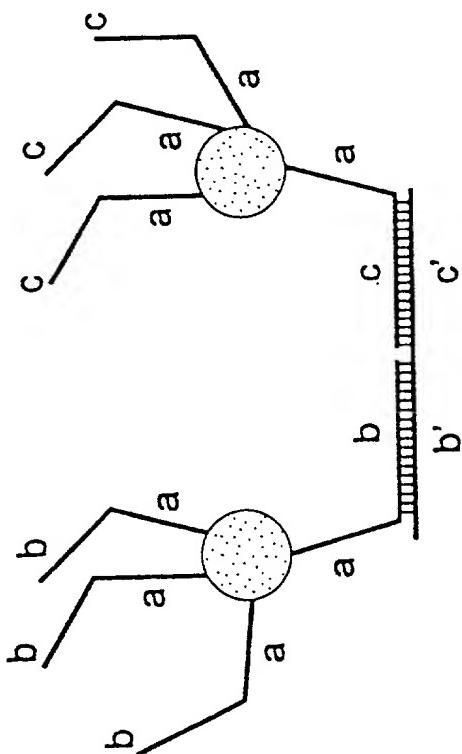


FIG. 17C

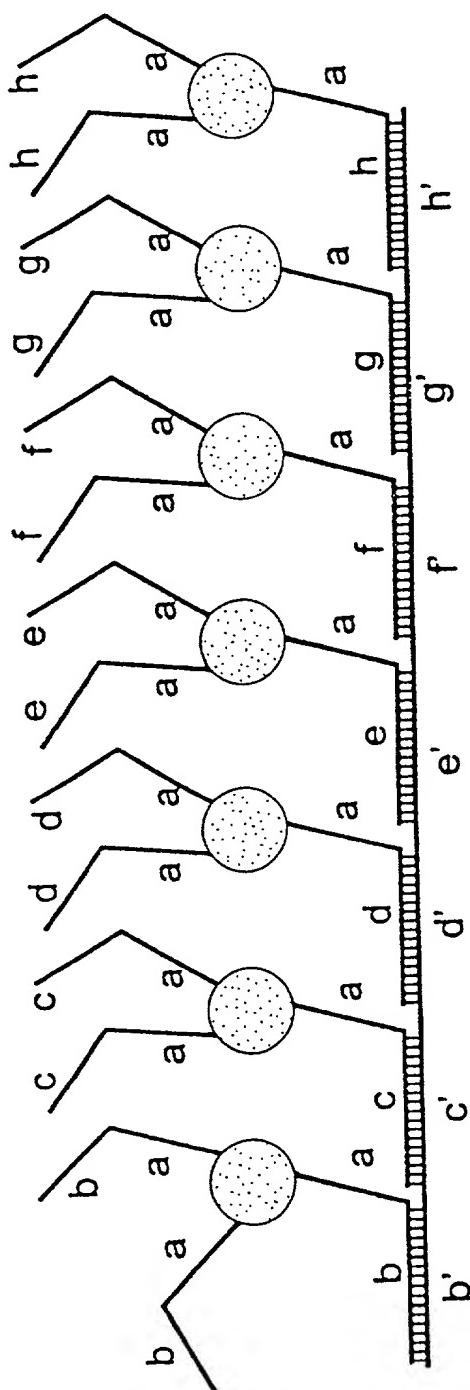


FIG. 17D

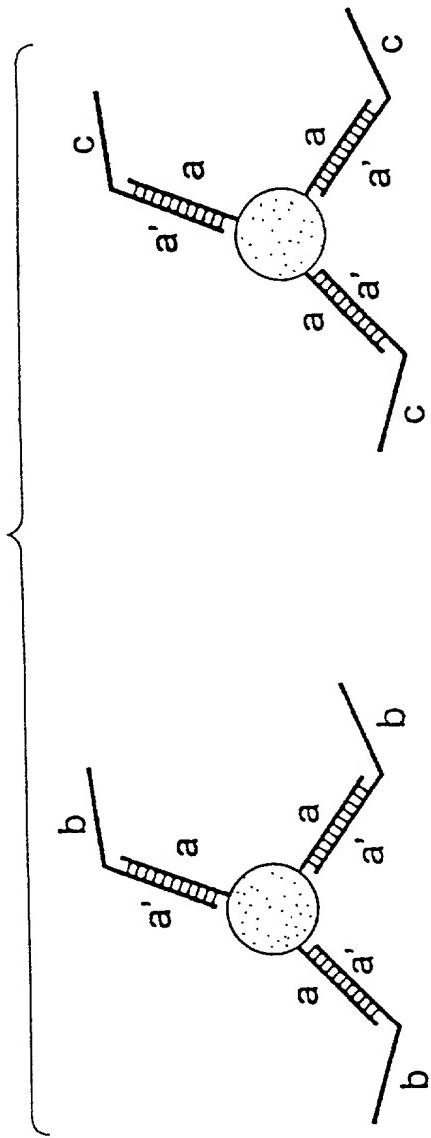


FIG. 17E

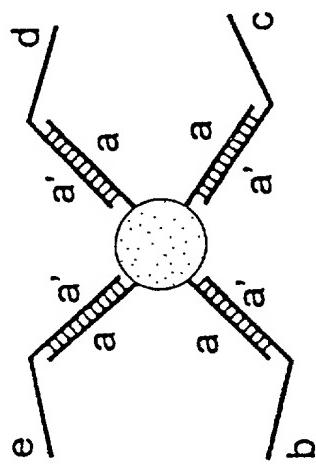


FIG. 18

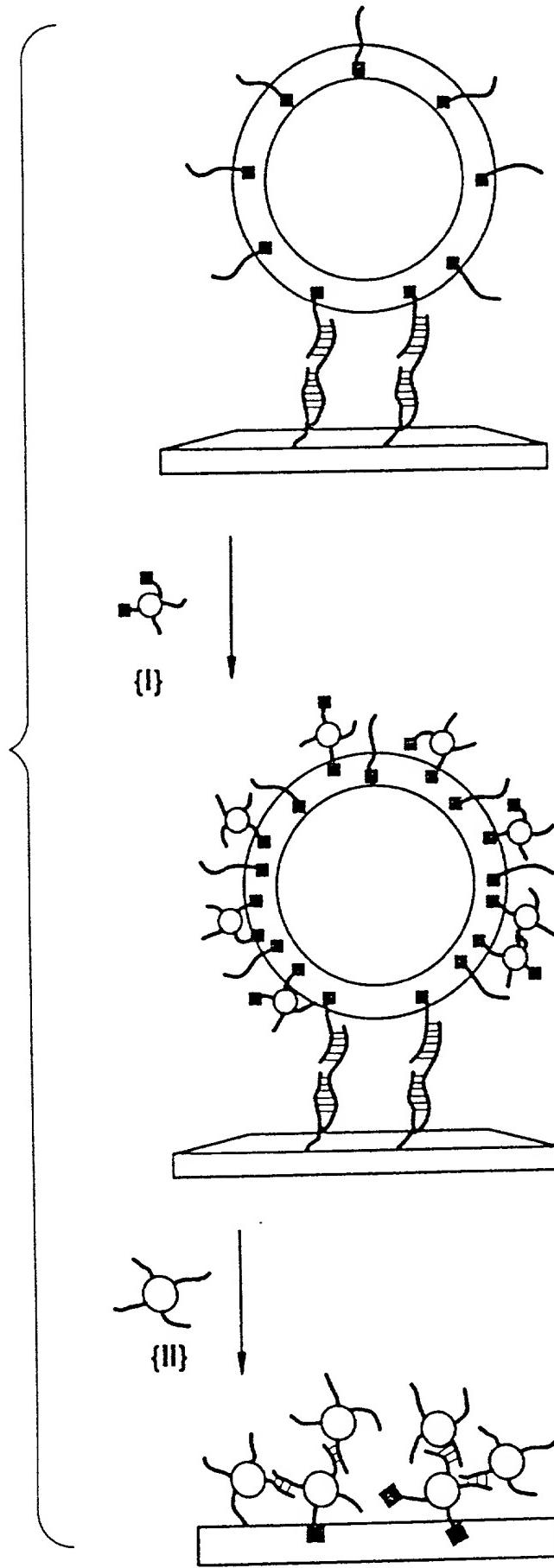


FIG. 19A

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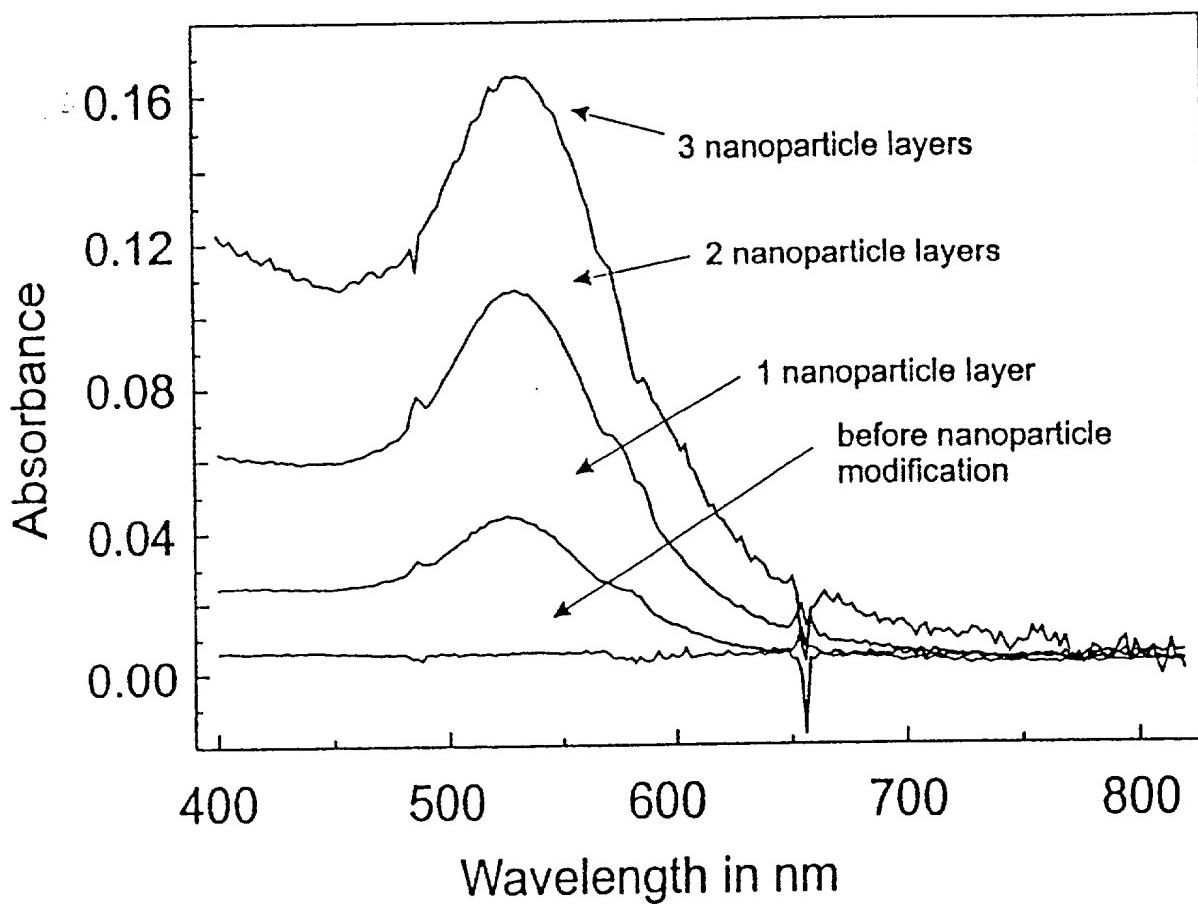


FIG. 19B

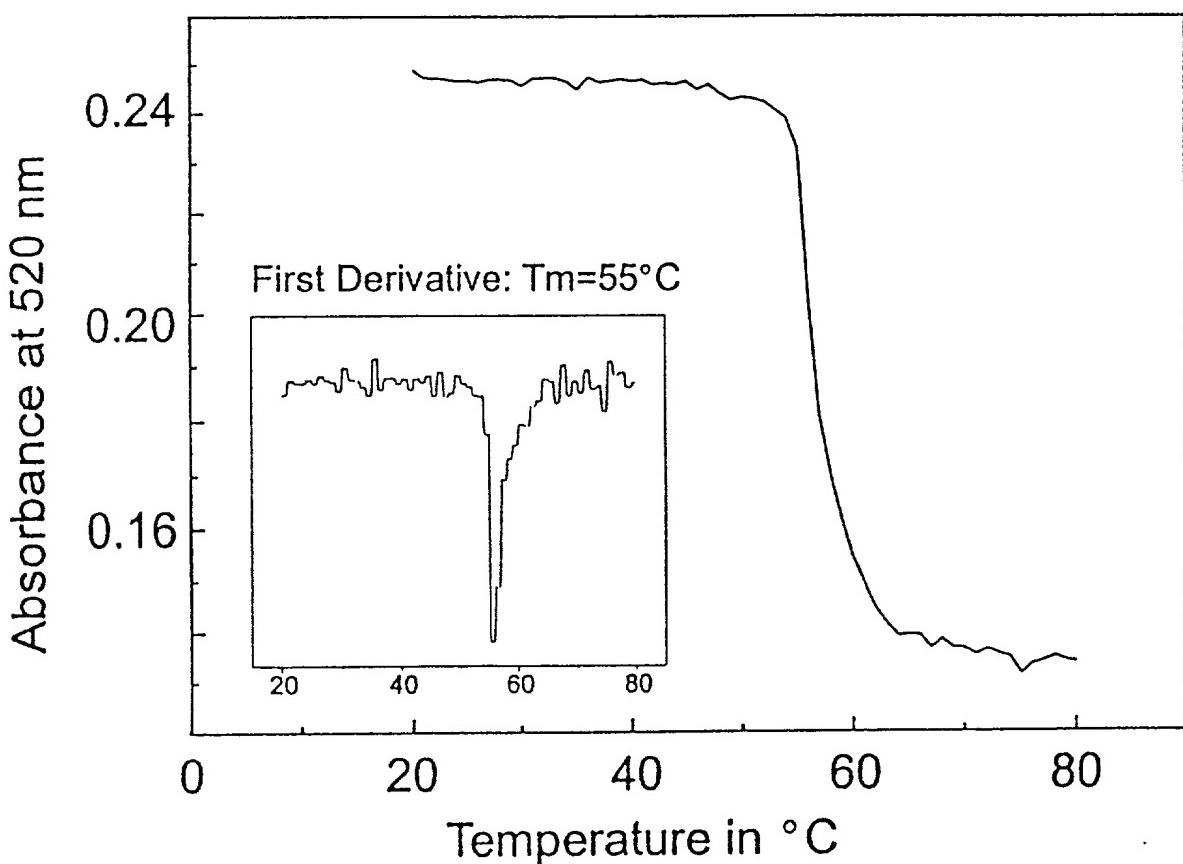


FIG.20A

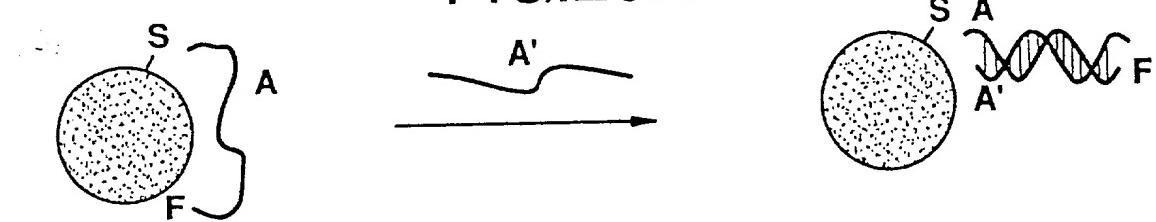
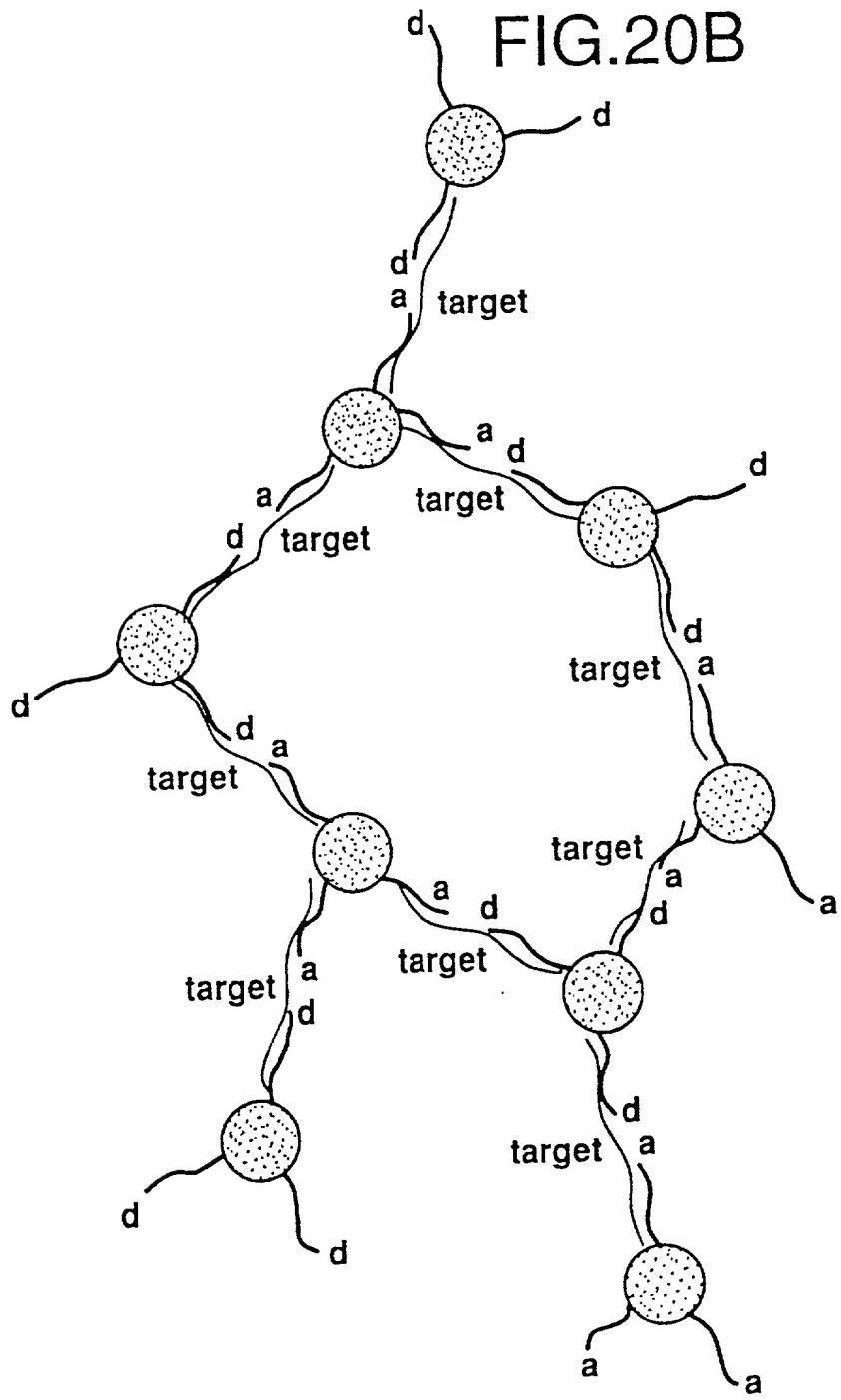


FIG.20B



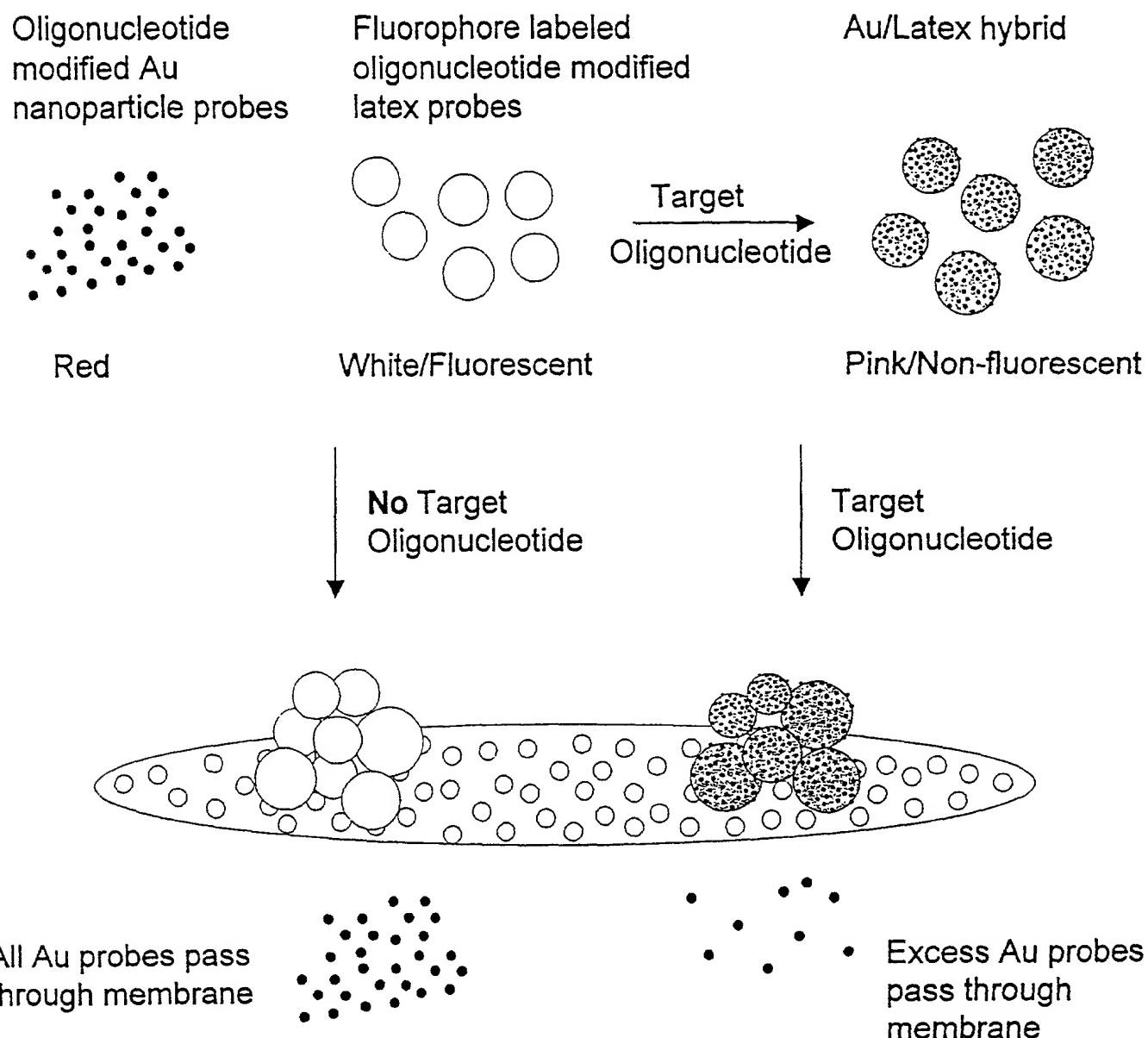
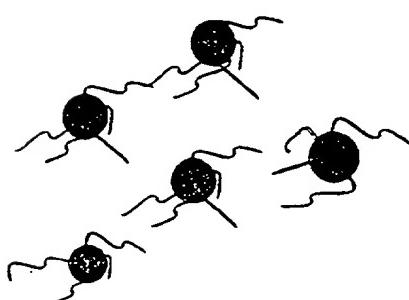


FIGURE 21

FIGURE 22

Fluorescent

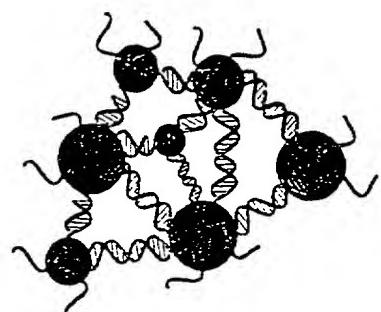
Nanoparticle Probes



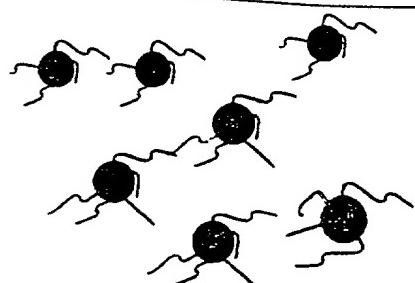
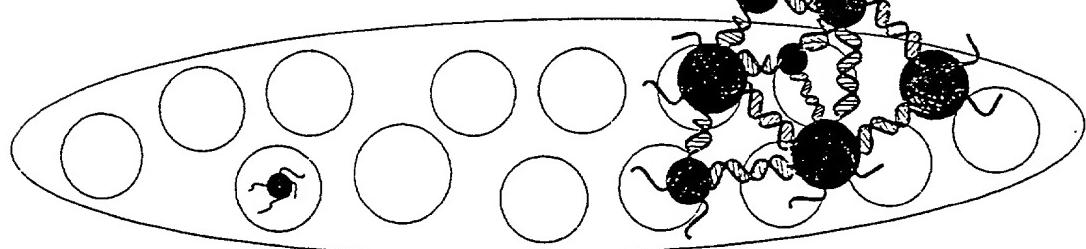
Target  
Oligonucleotide

Fluorescent

Cross-linked Aggregates



Target  
Oligonucleotide



The fluorescent nanoparticle probes  
pass through the membrane

The fluorescent cross-linked aggregates  
are retained by the membrane

## Anthrax PCR Product

5' G GCG GAT GAG TCA GTA GTT AAG GAG GCT CAT AGA GAA GTA ATT AAT  
3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT A TCT CTT CAT TAA TTA

TCG TCA ACA GAG GGA TTA TTG TTA AAT ATT GAT AAG GAT ATA AGA AAA  
AGC AGT TGT CTC CCT AAT AAC AAT TTA TAA CTA TTC CTA TAT TCT TTT

ATA TTA TCC AGG GTT ATA TTG TAG AAA TTG AAG ATA CTG AAG GGC TT 3'  
TAT AAT AGG TCC CAA TAT AAC ATC TTT AAC TTC TAT GAC TTC CCG AA 5'

141 mer Anthrax PCR product

[SEQ ID NO:36]

3' CTC CCT AAT AAC AAT  
[SEQ ID NO:37]

3' TTA TAA CTA TTC CTA  
[SEQ ID NO:38]

Oligonucleotide-Nanoparticle Probes

### Blocker Oligonucleotides

3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT

[SEQ ID NO:39]

3' A TCT CTT CAT TAA TTA AGC AGT TGT

[SEQ ID NO:40]

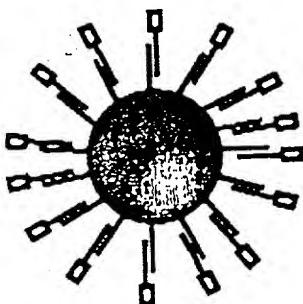
3' TAT TCT TTT TAT AAT AGG TCC CAA TAT

[SEQ ID NO:41]

3' AAC ATC TTT AAC TTC TAT GAC TTC CCG AA

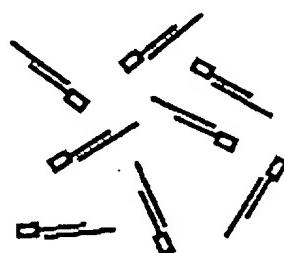
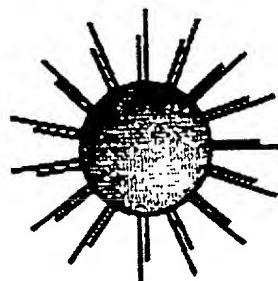
[SEQ ID NO:42]

FIGURE 23



Satellite Probe

Target  
→



Detection Signal

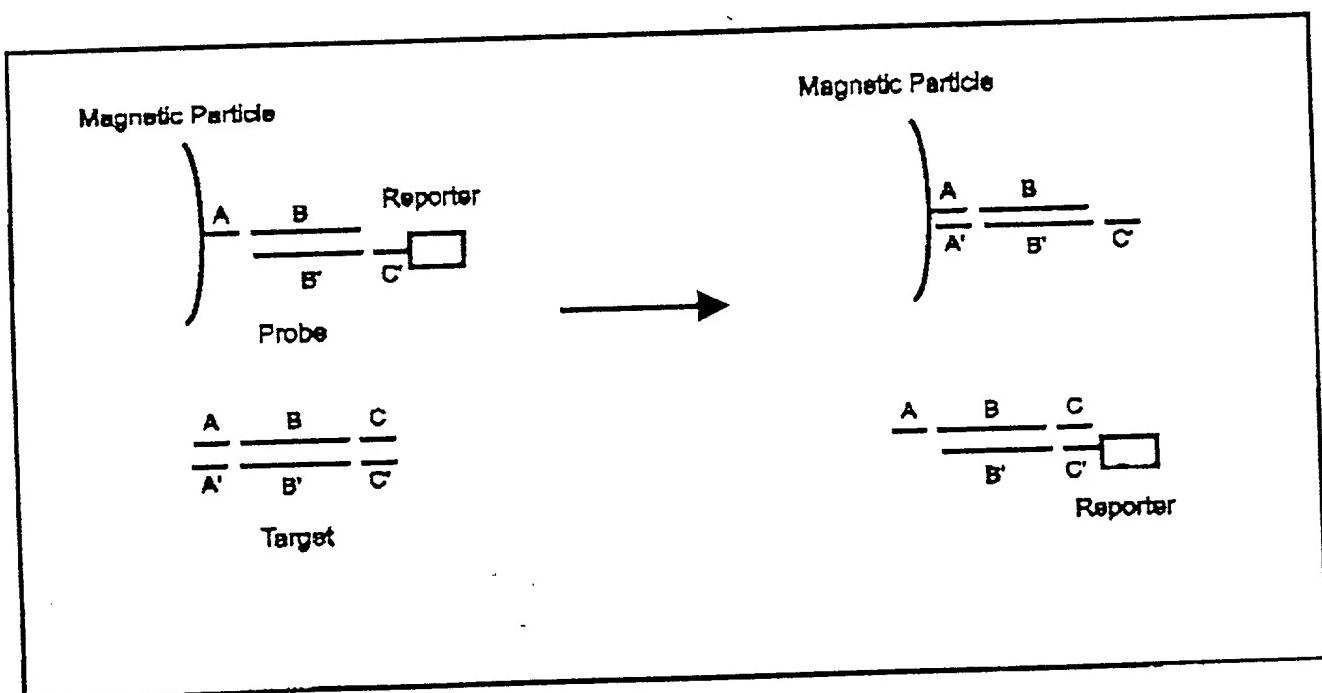


FIGURE 24